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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/756,690A

DATE: 05/02/2002 P.6
TIME: 16:44:01

Input Set : A:\249-124.app

Output Set: N:\CRF3\05022002\I756690A.raw

4 <110> APPLICANT: KOLTERMAN, ORVILLE G.
 5 YOUNG, ANDREW A.
 7 <120> TITLE OF INVENTION: USE OF EXENDINS AND AGONISTS THEREOF FOR MODULATION OF
 8 TRIGLYCERIDE LEVELS AND TREATMENT OF DYSLIPIDEMIA
 10 <130> FILE REFERENCE: 249/124
 12 <140> CURRENT APPLICATION NUMBER: 09/756,690A
 C--> 13 <141> CURRENT FILING DATE: 2002-04-19
 15 <150> PRIOR APPLICATION NUMBER: 60/175,365
 16 <151> PRIOR FILING DATE: 2000-01-10
 18 <160> NUMBER OF SEQ ID NOS: 188
 20 <170> SOFTWARE: PatentIn Ver 2.1
 22 <210> SEQ ID NO: 1
 23 <211> LENGTH: 39
 24 <212> TYPE: PRT
 25 <213> ORGANISM: Heloderma horridum
 27 <220> FEATURE:
 28 <223> OTHER INFORMATION: Exendin-3
 30 <220> FEATURE:
 31 <223> OTHER INFORMATION: c-term amidation
 33 <400> SEQUENCE: 1
 34 His Ser Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 35 1 5 10 15
 37 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 38 20 25 30
 40 Ser Gly Ala Pro Pro Pro Ser
 41 35
 44 <210> SEQ ID NO: 2
 45 <211> LENGTH: 39
 46 <212> TYPE: PRT
 47 <213> ORGANISM: Heloderma suspectum
 49 <220> FEATURE:
 50 <223> OTHER INFORMATION: Exendin-4
 52 <220> FEATURE:
 53 <223> OTHER INFORMATION: c-term amidation
 55 <400> SEQUENCE: 2
 56 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 57 1 5 10 15
 59 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 60 20 25 30
 62 Ser Gly Ala Pro Pro Pro Ser
 63 35
 66 <210> SEQ ID NO: 3
 67 <211> LENGTH: 39

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Input Set : A:\249-124.app

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68 <212> TYPE: PRT
69 <213> ORGANISM: Artificial Sequence
71 <220> FEATURE:
72 <223> OTHER INFORMATION: Description of Artificial Sequence: Exendin Agonist
74 <220> FEATURE:
75 <221> NAME/KEY: MOD_RES
76 <222> LOCATION: (1)
77 <223> OTHER INFORMATION: His, Arg or Tyr
79 <220> FEATURE:
80 <221> NAME/KEY: MOD_RES
81 <222> LOCATION: (2)
82 <223> OTHER INFORMATION: Ser, Gly, Ala or Thr
84 <220> FEATURE:
85 <221> NAME/KEY: MOD_RES
86 <222> LOCATION: (3)
87 <223> OTHER INFORMATION: Asp or Glu
89 <220> FEATURE:
90 <221> NAME/KEY: MOD_RES
91 <222> LOCATION: (6)
92 <223> OTHER INFORMATION: Phe, Tyr or naphthylalanine
94 <220> FEATURE:
95 <221> NAME/KEY: MOD_RES
96 <222> LOCATION: (7)
97 <223> OTHER INFORMATION: Thr or Ser
99 <220> FEATURE:
100 <221> NAME/KEY: MOD_RES
101 <222> LOCATION: (8)
102 <223> OTHER INFORMATION: Ser or Thr
104 <220> FEATURE:
105 <221> NAME/KEY: MOD_RES
106 <222> LOCATION: (9)
107 <223> OTHER INFORMATION: Asp or Glu
109 <220> FEATURE:
110 <221> NAME/KEY: MOD_RES
111 <222> LOCATION: (10)
112 <223> OTHER INFORMATION: Leu, Ile, Val, pentylglycine or Met
114 <220> FEATURE:
115 <221> NAME/KEY: MOD_RES
116 <222> LOCATION: (14)
117 <223> OTHER INFORMATION: Leu, Ile, pentylglycine, Val or Met
119 <220> FEATURE:
120 <221> NAME/KEY: MOD_RES
121 <222> LOCATION: (22)
122 <223> OTHER INFORMATION: Phe, Tyr or naphthylalanine
124 <220> FEATURE:
125 <221> NAME/KEY: MOD_RES
126 <222> LOCATION: (23)
127 <223> OTHER INFORMATION: Ile, Val, Leu, pentylglycine, tert-butylglycine or Met
129 <220> FEATURE:

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130 <221> NAME/KEY: MOD_RES
131 <222> LOCATION: (24)
132 <223> OTHER INFORMATION: Glu or Asp
134 <220> FEATURE:
135 <221> NAME/KEY: MOD_RES
136 <222> LOCATION: (25)
137 <223> OTHER INFORMATION: Trp, Phe, Tyr, or naphthylalanine
139 <220> FEATURE:
140 <221> NAME/KEY: MOD_RES
141 <222> LOCATION: (31)
142 <223> OTHER INFORMATION: Pro, homoproline, 3Hyp, 4Hyp, thioproline,
143     N-alkylglycine, N-alkylpentylglycine, or N-alkylalanine
145 <220> FEATURE:
146 <221> NAME/KEY: MOD_RES
147 <222> LOCATION: (36)
148 <223> OTHER INFORMATION: Pro, homoproline, 3Hyp, 4Hyp, thioproline,
149     N-alkylglycine, N-alkylpentylglycine, or N-alkylalanine
151 <220> FEATURE:
152 <221> NAME/KEY: MOD_RES
153 <222> LOCATION: (37)
154 <223> OTHER INFORMATION: Pro, homoproline, 3Hyp, 4Hyp, thioproline,
155     N-alkylglycine, N-alkylpentylglycine, or N-alkylalanine
157 <220> FEATURE:
158 <221> NAME/KEY: MOD_RES
159 <222> LOCATION: (38)
160 <223> OTHER INFORMATION: Pro, homoproline, 3Hyp, 4Hyp, thioproline,
161     N-alkylglycine, N-alkylpentylglycine, or N-alkylalanine
163 <220> FEATURE:
164 <221> NAME/KEY: MOD_RES
165 <222> LOCATION: (39)
166 <223> OTHER INFORMATION: Ser, Thr or Tyr, which is optionally amidated with the
167     provision that the compound is not exendin-3 or exendin-4
169 <400> SEQUENCE: 3
W--> 170 Xaa Xaa Xaa Gly Thr Xaa Xaa Xaa Xaa Ser Lys Gln Xaa Glu Glu
      171 1           5           10           15
W--> 173 Glu Ala Val Arg Leu Xaa Xaa Xaa Xaa Leu Lys Asn Gly Gly Xaa Ser
      174           20           25           30
W--> 176 Ser Gly Ala Xaa Xaa Xaa Xaa
      177           35
180 <210> SEQ ID NO: 4
181 <211> LENGTH: 38
182 <212> TYPE: PRT
183 <213> ORGANISM: Artificial Sequence
185 <220> FEATURE:
186 <223> OTHER INFORMATION: Description of Artificial Sequence: Exendin Agonist
188 <220> FEATURE:
189 <221> NAME/KEY: MOD_RES
190 <222> LOCATION: (1)
191 <223> OTHER INFORMATION: His, Arg or Tyr

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Input Set : A:\249-124.app

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193 <220> FEATURE:
194 <221> NAME/KEY: MOD_RES
195 <222> LOCATION: (2)
196 <223> OTHER INFORMATION: Ser, Gly, Ala or Thr
198 <220> FEATURE:
199 <221> NAME/KEY: MOD_RES
200 <222> LOCATION: (3)
201 <223> OTHER INFORMATION: Asp or Glu
203 <220> FEATURE:
204 <221> NAME/KEY: MOD_RES
205 <222> LOCATION: (5)
206 <223> OTHER INFORMATION: Ala or Thr
208 <220> FEATURE:
209 <221> NAME/KEY: MOD_RES
210 <222> LOCATION: (6)
211 <223> OTHER INFORMATION: Ala, Phe, Tyr or naphthylalanine
213 <220> FEATURE:
214 <221> NAME/KEY: MOD_RES
215 <222> LOCATION: (7)
216 <223> OTHER INFORMATION: Thr or Ser
218 <220> FEATURE:
219 <221> NAME/KEY: MOD_RES
220 <222> LOCATION: (8)
221 <223> OTHER INFORMATION: Ala, Ser or Thr
223 <220> FEATURE:
224 <221> NAME/KEY: MOD_RES
225 <222> LOCATION: (9)
226 <223> OTHER INFORMATION: Asp or Glu
228 <220> FEATURE:
229 <221> NAME/KEY: MOD_RES
230 <222> LOCATION: (10)
231 <223> OTHER INFORMATION: Ala, Leu, Ile, Val, pentylglycine or Met
233 <220> FEATURE:
234 <221> NAME/KEY: MOD_RES
235 <222> LOCATION: (11)
236 <223> OTHER INFORMATION: Ala or Ser
238 <220> FEATURE:
239 <221> NAME/KEY: MOD_RES
240 <222> LOCATION: (12)
241 <223> OTHER INFORMATION: Ala or Lys
243 <220> FEATURE:
244 <221> NAME/KEY: MOD_RES
245 <222> LOCATION: (13)
246 <223> OTHER INFORMATION: Ala or Gln
248 <220> FEATURE:
249 <221> NAME/KEY: MOD_RES
250 <222> LOCATION: (14)
251 <223> OTHER INFORMATION: Ala, Leu, Ile, pentylglycine, Val or Met
253 <220> FEATURE:

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Input Set : A:\249-124.app

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254 <221> NAME/KEY: MOD_RES
255 <222> LOCATION: (15)
256 <223> OTHER INFORMATION: Ala or Glu
258 <220> FEATURE:
259 <221> NAME/KEY: MOD_RES
260 <222> LOCATION: (16)
261 <223> OTHER INFORMATION: Ala or Glu
263 <220> FEATURE:
264 <221> NAME/KEY: MOD_RES
265 <222> LOCATION: (17)
266 <223> OTHER INFORMATION: Ala or Glu
268 <220> FEATURE:
269 <221> NAME/KEY: MOD_RES
270 <222> LOCATION: (19)
271 <223> OTHER INFORMATION: Ala or Val
273 <220> FEATURE:
274 <221> NAME/KEY: MOD_RES
275 <222> LOCATION: (20)
276 <223> OTHER INFORMATION: Ala or Arg
278 <220> FEATURE:
279 <221> NAME/KEY: MOD_RES
280 <222> LOCATION: (21)
281 <223> OTHER INFORMATION: Ala or Leu
283 <220> FEATURE:
284 <221> NAME/KEY: MOD_RES
285 <222> LOCATION: (22)
286 <223> OTHER INFORMATION: Ala, Phe, Tyr or naphthylalanine
288 <220> FEATURE:
289 <221> NAME/KEY: MOD_RES
290 <222> LOCATION: (23)
291 <223> OTHER INFORMATION: Ile, Val, Leu, pentylglycine, tert-butylglycine or Met
293 <220> FEATURE:
294 <221> NAME/KEY: MOD_RES
295 <222> LOCATION: (24)
296 <223> OTHER INFORMATION: Ala, Glu or Asp
298 <220> FEATURE:
299 <221> NAME/KEY: MOD_RES
300 <222> LOCATION: (25)
301 <223> OTHER INFORMATION: Ala, Trp, Phe, Tyr or naphthylalanine
303 <220> FEATURE:
304 <221> NAME/KEY: MOD_RES
305 <222> LOCATION: (26)
306 <223> OTHER INFORMATION: Ala or Leu
308 <220> FEATURE:
309 <221> NAME/KEY: MOD_RES
310 <222> LOCATION: (27)
311 <223> OTHER INFORMATION: Ala or Lys
313 <220> FEATURE:
314 <221> NAME/KEY: MOD_RES

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/756,690A

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Input Set : A:\249-124.app
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 1,2,3,6,7,8,9,10,14,22,23,24,25,31,36,37,38,39
Seq#:4; Xaa Pos. 1,2,3,5,6,7,8,9,10,11,12,13,14,15,16,17,19,20,21,22,23,24
Seq#:4; Xaa Pos. 25,26,27,28,31,36,37,38
Seq#:5; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,19,20,21,22,23
Seq#:5; Xaa Pos. 24,25,26,27,28,31,36,37,38
Seq#:15; Xaa Pos. 6
Seq#:20; Xaa Pos. 10
Seq#:21; Xaa Pos. 10
Seq#:22; Xaa Pos. 14
Seq#:23; Xaa Pos. 14
Seq#:24; Xaa Pos. 22
Seq#:27; Xaa Pos. 23
Seq#:28; Xaa Pos. 23
Seq#:31; Xaa Pos. 31,36,37,38
Seq#:32; Xaa Pos. 36,37,38
Seq#:33; Xaa Pos. 31,36,37,38
Seq#:34; Xaa Pos. 36,37,38
Seq#:35; Xaa Pos. 31,36,37,38
Seq#:36; Xaa Pos. 31,36,37,38
Seq#:37; Xaa Pos. 31,36,37,38
Seq#:38; Xaa Pos. 36,37,38
Seq#:39; Xaa Pos. 31,36,37,38
Seq#:81; Xaa Pos. 31,36,37,38
Seq#:82; Xaa Pos. 36,37,38
Seq#:83; Xaa Pos. 31
Seq#:84; Xaa Pos. 31,36,37
Seq#:85; Xaa Pos. 31,36,37
Seq#:86; Xaa Pos. 31,36
Seq#:89; Xaa Pos. 6
Seq#:93; Xaa Pos. 10
Seq#:94; Xaa Pos. 22
Seq#:95; Xaa Pos. 23
Seq#:99; Xaa Pos. 31,36,37
Seq#:115; Xaa Pos. 6
Seq#:116; Xaa Pos. 6
Seq#:127; Xaa Pos. 10
Seq#:128; Xaa Pos. 10
Seq#:137; Xaa Pos. 14
Seq#:138; Xaa Pos. 14
Seq#:151; Xaa Pos. 22
Seq#:152; Xaa Pos. 22
Seq#:155; Xaa Pos. 23
Seq#:156; Xaa Pos. 23
Seq#:181; Xaa Pos. 31,36,37,38

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Seq#:182; Xaa Pos. 36,37,38

Seq#:183; Xaa Pos. 31,36,37

Seq#:184; Xaa Pos. 31,36

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Input Set : A:\249-124.app

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:170 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:173 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:16
L:176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:32
L:345 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:348 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:16
L:351 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:32
L:530 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:533 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:16
L:536 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:32
L:742 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:857 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:884 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:911 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:938 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:968 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:16
L:1039 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:16
L:1066 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:16
L:1152 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:16
L:1155 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:32
L:1192 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:32
L:1231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:16
L:1234 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:32
L:1271 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:32
L:1310 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:16
L:1313 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:32
L:1352 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:16
L:1355 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:32
L:1394 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:16
L:1397 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:32
L:1434 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:32
L:1473 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:16
L:1476 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:32
L:2325 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81 after pos.:16
L:2328 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81 after pos.:32
L:2365 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82 after pos.:32
L:2389 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83 after pos.:16
L:2426 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84 after pos.:16
L:2429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84 after pos.:32
L:2463 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85 after pos.:16
L:2466 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85 after pos.:32
L:2495 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86 after pos.:16
L:2498 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86 after pos.:32
L:2560 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:89 after pos.:0
L:2641 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93 after pos.:0
L:2668 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94 after pos.:16
L:2692 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:95 after pos.:16
L:2785 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:99 after pos.:16

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L:2788 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:99 after pos.:32
L:3093 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:115 after pos.:0
L:3117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:116 after pos.:0